226/284 Patent

CLAIMS

What we claim is:

1. A plasmid for expression of recombinant eucaryotic genes comprising:

a first transcription unit comprising a first transcriptional control sequence transcriptionally linked with a first 5'-untranslated region, a first synthetic intron, a first coding sequence, and a first synthetic 3'-untranslated region/poly(A) signal, wherein said first synthetic intron is between said control sequence and said first coding sequence; and

a second transcription unit comprising a second transcriptional control sequence transcriptionally linked with a second 5'-untranslated region, a second synthetic intron, a second coding sequence, and a second synthetic 3'-untranslated region/poly(A) signal, wherein said second synthetic intron is between said control sequence and said second coding sequence.

- 2. The plasmid of claim 1, wherein said first transcriptional control sequence or said second transcriptional control sequence comprise cytomegalovirus promoter/enhancer sequences.
- 3. The plasmid of claim 1, wherein said first coding sequence or said second coding sequence encode a therapeutic molecule or a subunit of a therapeutic molecule.

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- 4. The plasmid of claim 1, wherein said first and second transcriptional control sequences are the same.
- 5. The plasmid of claim 1, wherein said first and second transcriptional control sequences are different.
- 6. The plasmid of claim 1, wherein said first coding sequence and said second coding sequence comprise a sequence coding for the p40 subunit of human IL-12 and a sequence coding for the p85 subunit of human IL-12.
- 7. The plasmid of claim 6, wherein said sequence coding for the p40 subunit of human IL-12 is 5' to said sequence coding for the p35 subunit of human IL-12.
- A plasmid for expression of recombinant eucaryotic genes, comprising an intron having variable splicing, a first coding sequence, and a second coding sequence.
- 9. The plasmid of claim 8, further comprising:
 a transcriptional control sequence
 transcriptionally linked with a first coding sequence
 and a second coding sequence;
 - a 5'-untranslated region;

an intron 5' to said first coding sequence; an alternative splice site 3' to said first coding sequence and 5' to said second coding sequence; and

a 3'-untranslated region/poly(A) signal.

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10. The plasmid of claim 9, wherein said first coding sequence or said second coding sequence encode a therapeutic molecule or a subunit of a therapeutic molecule.

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11. The plasmid of claim 9, wherein said transcriptional control sequence comprises a cytomegalovirus promoter enhancer sequence.

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12. The plasmid of claim 8, wherein said first coding sequence and said second coding sequence comprise a sequence coding for the p40 subunit of human IL-12 and a sequence coding for the p35 subunit of human IL-12.

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13. A plasmid for expression of recombinant eucaryotic genes comprising:

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a transcriptional control sequence transcriptionally linked with a first coding sequence, an IRES sequence, a second coding sequence, and a 3'untranslated region/poly(A) signal, wherein said IRES sequence is between said first coding sequence and said second coding sequence; and

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an intron between said promoter and said first coding sequence.

14. The plasmid of claim 13, wherein said transcriptional control sequence comprises a cytomegalovirus promoter/enhancer sequence.

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- 15. The plasmid of claim 13, wherein said first coding sequence and said second coding sequence comprise a sequence coding for the p40 subunit of human IL-12 and a sequence coding for the p35 subunit of human IL-12.
- 16. The plasmid of claim 13, wherein said IRES sequence is from an encephalomy carditis virus.
- A DNA sequence coding for human IL-12 subunit, comprising a synthetic nucleotide sequence having less than 50% identity to a matural human IL-12 subunit coding sequence.
- 18. The DNA sequence of claim 17, wherein said synthetic nucleotide sequence comprises a sequence having at least 99% sequence identity to the sequence of SEQ ID NO. 3.
- 19. The DNA sequence of claim 18, wherein said synthetic nucleotide sequence comprises a nucleotide sequence identical to the sequence of SEQ ID NO. 3 or 4.
- 20. The DNA sequence of claim 17, wherein said synthetic nucleotide sequence comprises a sequence having at least 99% sequence identity to the sequence of SEQ ID NO. 7.
- 21. The DNA sequence of claim 20, wherein said synthetic nucleotide sequence comprises a nucleotide

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28. The composition of claim 27, wherein said isotonic carbohydrate solution consists essentially of about 10% lactose.

- 22. A composition for delivery of a DNA molecule in a mammal, comprising
- a cationic lipid with a neutral co-lipid, prepared as a liposome having an extrusion size of about 800 nanometers; and
 - a quantity of DNA comprising a coding sequence.
- 23. The composition of claim 22, wherein said DNA is at least about 80% supercoiled.
- 24. The composition of claim 23, wherein said DNA is at least about 90% supercolled.
- 25. The composition of claim 24, wherein said DNA is at least about 95% supercoiled.
- 26. The composition of claim 22, wherein said cationic lipid and said DNA are present in a negative to positive charge ratio of about 1:3.
- 27. The composition of claim 22, further comprising an isotonic carbohydrate solution.

- 29. A composition of claim 22, wherein said cationic lipid is DOTMA and said neutral co-lipid is cholesterol.
- 30. A composition for delivery of a DNA molecule in a mammal, comprising

a cationic lipid with a neutral co-lipid; and
a quantity of DNA comprising a coding sequence,
wherein said cationic lipid and said DNA are
present in a negative to positive charge ratio of about
1:3.

- 31. The composition of claim 30, wherein said DNA is at least about 80% supercoiled.
- 32. The composition of claim 31, wherein said DNA is at least about 90% supercoiled.
- 33. The composition of claim 32, wherein said DNA is at least about 95% supercoiled.
- 34. The composition of claim 30, further comprising an isotonic carbohydrate solution.
- 35. The composition of claim 34, wherein said isotonic carbohydrate solution consists essentially of about 10% lactose.
- 36. A composition of claim 30, wherein said SSSD/62166. v01

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cationic lipid is DOTMA and said neutral co-lipid is cholesterol.

A method for preparing a composition for delivery of a DNA to a mammal, comprising the steps of:

- preparing a DNA comprising a coding sequence;
- preparing liposomes having an extrusion size of about 800 nm, wherein said liposomes comprise a cationic lipid and a neutral co-lipid; and
- combining said liposomes with said DNA in amounts such that said cathonic lipid and said DNA are present in a negative to positive charge ratio of about 1:3.

A method of treatment of a mammalian condition or disease, comprising administering to a mammal suffering from said condition or disease an amount of a composition for delivery of a DNA molecule in a mammal,

wherein said DNA comprises a coding sequence encoding a therapeutic molecule or a subunit thereof, and

wherein said composition comprises a cationic lipid, a neutral co-lipid, and said DNA, and has a negative to positive charge ratio of about 1:3 for said cationic lipid and said DNA.

39. The method of claim 38,\wherein said composition is prepared for administration by ultrasonic nebulization.

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comprises	two	coding	se	quences	wh:	ich	encode	e hum	nan	IL-12
p40 and p3	35 sı	ubunits.	. \							

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- The method of claim 38, wherein said disease 41. or condition is asthma.
- The method of claim 38, wherein said disease 42. or condition is a cancer
- A vaccine adjuvant comprising a cationic lipid, a neutral co-lipid and DNA,

wherein said DNA/ comprises a sequence encoding the p40 subunit of IL-12/and the p35 subunit of IL-12, and

wherein said cationic lipid and said DNA are present in a negative to positive charge ratio of about 1:3.

A method of enhancing the response of a mammal to a vaccine, comprising the step of administering to said mammal a vaccine and an adjuvant,

wherein said adjuvant comprises a cationic lipid, a neutral co-lipid, and DNA, said DNA comprising a sequence encoding the p40 subunit of IL-12 and the p35 subunit of IL-12, and

wherein said cationic lipid and said DNA are present in a negative to positive charge ratio of about 1:3.

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